

Fig. 1A
SEQ. ID NO:1

MTVARPSVPAALPLLGLLPRLLLLVLCLPAVWGDCGLPPDVPNAQPALE 50
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PTRLNSASLKPQYITQNYFPVGTVEYECRPGYRREPSLSPKLTCLQNLK 150
WSTAVEFCKKKSCPNPGEIRNGQIDVPGGILFGATISFSCNTGYKLFGST 200
SSFCLISGSSVQWSDDLPECREIYCPAPPQIDNGIIQGERDHYGYRQSVT 250
YACNKGFTMIGEHSIYCTVNNDGEWSGPPPECRGKSLTSKVPPTVQKPT 300
TVNVPTEVSPTSQKTTTKTTTPNAQATRSTPVSRTTKHFHETTPNKGSG 350
TTSGETRLLSGHTCFTLTGLLGLTMTGLLT

Fig. 1B
SEQ. ID NO:2

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1981 atttatattt atttatgaca gtgaacattc tgattttaca tgtaaaacaa gaaaagttga
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Fig. 2

SEQ. ID NO:3

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WSPAAPTCEVKSCDDFMGQLLNGRVLPVNLQLGAKVDFVCDEGFQLKGS 400
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Fig. 3
SEQ. ID NO:4

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Fig. 4A

SEQ. ID NO:5

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TVMFECDKG FYLDGSDTIVCDNSNSTWDPPVPKCLKVSTSTTKSPASSAS 300
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Fig. 4B

SEQ. ID NO:6

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```

Fig. 5

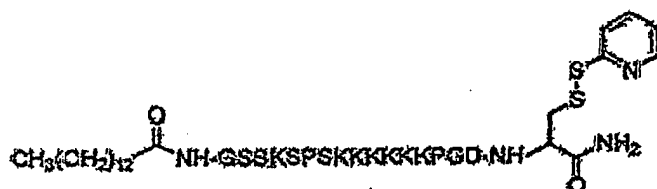


Fig. 6A

SEQ. ID NO:7

ATA TAC GAA TTC AGA TCT ATG ACC GTC GCG CGG CCG AGC GTG

Fig. 6B

SEQ. ID NO:8

ACA GTG CTC GAG CAT TCA GGT GGT GGG CCA CTC CA

Fig 7A

SEQ. ID NO:9

ATA TAC CTC GAG TCC TAA CAA ATG CAC GCC TCC AAA TGT GG-3

Fig 7B

SEQ. ID NO:10

ACA GTG ATG CAT TGG TTT GGG TTT TCA ACT TGG C

Fig 7C

SEQ. ID NO:11

ATA TAC ATG CAT CTG ACT TTC CCA TTG GGA CAT CTT TAA AG

Fig 7D

SEQ. ID NO:12

ACA GTG AGA TCT TTA GTG ATG GTG ATG GTG ATG AAT TCC ACA GCG AGG GGC
AGG GCT

Fig. 8A
SEQ ID NO:13

M T V A R P S V P A A L P
L L G E L P P R L L L L V L L C L P A V W G
D C G L P P D V P N A Q P A L E G R T S F
P E D T V I T Y K C E E S F V K I P G E K
D S V I C L K G S Q W S D I E E F C N R S
C E V P T R L N S A S L K Q P G Y I T Q N Y
F P V G T V L N E A C R P G Y R E F C K K
S S C L P N P G E I R N G Q I D V P G G I L
F G A T I S S S C N T G Y K L F G S T S S
Y C P A P P Q I D N G I I Q G E R D H Y G
Y R Q S V T Y A C N K G F T M I G E H S I
Y C T V N N D E G E W S G P P P E C
S S P N K C T P P N V E N G I L V S D N
R S L F S L N E V V E F R C Q P G F V M
K G P R R V K C Q Q A L N K W E P E L P S
C S R V C Q P P D V L H A E E R T Q R D
K D N F S P G Q E V F Y S C E P G Y D L
R G A A S M R C T P Q Q G D W S P A V P T
C E V K S C D D F M G Q L L N G E G F Q L
P V N L Q L G A K V D F V C D E G F Q L
K G S S A S Y C V L A G M E S L W N S S
V P V C E Q I F C P S P P V I P N G R H
T G K P L E V F P F G K A V N Y T C D P
H P D R G T S F D L I G E S T I R C T S
D P Q G N G V W S S P A P R C G I L G H
C Q A P D H F L F A K L K T Q T N A S D
F P I G T S L K Y E C R P E Y Y G R P F
S I T C L D N L V W S S P K D V C K R K
S C K T P P D P V N G M V H V I T D I Q
V G S R I N Y S C T T G H R L I G H S S
A E C I L S G N A A H W S T K P P I C Q
R I P C G L P P T I A N G D F I S T N R
E N F H Y G S V V T Y R C N P G S G R
K V F E L V G P A P Q S I I P N K C T P P N
G I W S G I L V S D N R S L F S L N E V V
V E N G I P G F V M K G P R R V K C Q A
E F R C Q P E L P S C S R V C Q P P P D
L N K W E P E L P S C S R V C Q P P P D
V L H A E R T Q R D K D N F S P G Q E V
F Y S C E P G Y D L R G A A S M R C T P
Q G D W S P A A P T C E V K S C D D F M
G Q L L N G R V L F P V N L Q L G A K V
D F V C D E G F Q L K G S S A S Y C V L
A G M E S L W N G S R H T G K P L E V F P
S P P V I P N G R H T G K P L E V F P
G K A V N Y T C D P H P D R G T S F D L

I G E S T I R C T S D P Q G N G V W S S
P A P R C G I H H H H H H

Fig. 8B

SEQ. ID NO: 14

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CTCCTCGGGGAGCTGCCCCGGCTGCTGCTGCTGGTGCTGTTGTGCTGCTGCCCCGGCTGTTGGGGT
GACTGTGGCCTTCCCCAGATGTACCTAATGCCAGCCAGCTTTGGAAGGCCGTACAAGTTT
CCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTTGTGAAAATTCCTGGCGAGAAG
GACTCAGTGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
TGCGAGGTGCCAACAAAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCAGGTTACAGAAGAGAACCTTCTCTA
TCACCAAACTAACTTGCCTTCAGAATTTAAAAATGGTCCACAGCAGTCGAATTTGTAAAAAG
AAATCATGCCCTAATCCGGGAGAAATACGAAATGGTCAGATTGATGTACCAGGTGGCATATTA
TTTGGTGCAACCATCTCCTTCTCATGTAACACAGGGTACAAATTATTTGGCTCGACTTCTAGT
TTTTGTCTTATTTAGGCAGCTCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAGAAATT
TATTGTCCAGCACCACCACAAATTGACAATGGAATAATTCAAGGGGAACGTGACCATTATGGA
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TATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCCACCACCTGAATGC
TCGAGTCCTAACAAATGCACGCCTCCAAATGTGGAATAATGGAATAATTGGTATCTGACAAC
AGAAGCTTATTTTCTTAAATGAAGTTGTGGAGTTTAGGTGTGAGCCTGGCTTTGTGTCATG
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TGCTCCAGGGTATGTGAGCCACCTCCAGATGTCTGTCATGCTGAGCGTACCCAAAGGGAC
AAGGACAACTTTTACCTGGGCAGGAAGTGTCTACAGCTGTGAGCCCGCTACGACCTC
AGAGGGGCTCGCTCTATGCGCTGCACACCCAGGGAGACTGGAGCCCTGCAGCCCCCACA
TGTAAGTGAATCCTGTGATGACTTCATGGGCCAACTTCTTAATGGCCGTGTGCTATTT
CCAGTAAATCTCAGCTTGGAGCAAAAGTGGATTTTTGTGTTGTGATGAAGGATTTCAATTA
AAAGGCAGCTCTGCTAGTTACTGTGTCTTGGCTGGAATGGAAAGCCTTTGGAATAGCAGT
GTTCCAGTGTGTGAACAAATCTTTTGTCCAAGTCCCTCCAGTTATTCTAATGGGAGACAC
ACAGGAAAACCTCTGGAAGTCTTTCCCTTTGGAAAAGCAGTAAATTACACATGCGACCCC
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TGTCAGCCCCAGATCATTTTCTGTTTGCCAAGTTGAAAACCCAAACCAATGCATCTGAC
TTTCCCATTTGGGACATCTTTAAAGTACGAATGCCGTCTGAGTACTACGGGAGGCCATT
TCTATCACATGTCTAGATAACCTGGTCTGGTCAAGTCCCAAAGATGTCTGTAAACGTAAA
TCATGTAAACTCCTCCAGATCCAGTGAATGGCATGGTGATGTGATCACAGACATCCAG
GTTGGATCCAGAATCAACTATCTTGTACTACAGGGCACCAGCTCATTGGTCACTCATCT
GCTGAATGTATCCTCTCGGGCAATGCTGCCCATTGGAGCACGAAGCCGCAATTTGTCAA
CGAATTCCTTGTGGGCTACCCCCACCATCGCCAATGGAGATTTTATTAGCACCACAGAG
GAGAATTTTCACTATGGATCAGTGGTGACCTACCGCTGCAATCCTGGAAGCGGAGGGAGA
AAGGTGTTTGTAGCTTGTGGGTGAGCCCTCCATATACTGCACCAGCAATGACGATCAAGTG
GGCATCTGGAGCGGCGCCCGGCCCTCAGTGCAATTATACCTAACAAATGCACGCCTCCAAAT
GTGGAATAATGGAATATTGGTATCTGACAACAGAAGCTTATTTTCTTAAATGAAGTTGTG
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GGCCAACCTTCTTAATGGCCGTGTGCTATTTCCAGTAAATCTCCAGCTTGGAGCAAAAGTG

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 AGTCCTCCAGTTATTCTTAAATGGGAGACACACAGGAAAACCTCTGGAAGTCTTTCCCTTT
 GGAAAAGCAGTAAATTACACATGCGACCCCCACCCAGACAGAGGGACGAGCTTCGACCTC
 ATTGGAGAGAGCACCATCCGCTGCACAAGTGACCCTCAAGGGAATGGGGTTTGGAGCAGC
 CCTGCCCCCTCGCTGTGGAATTCATCACCATCACCATCACTAAAGATCT

Fig. 9A

SEQ ID NO:15

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 M T V A R P S V P A A L P
 D C G L P P D V P N A Q P A L E G R T S F
 P E D T V I T Y K C E E S F V K I P G E K
 D S V I C L K G S Q W S D I E E F C N R S
 C E V P T R L N S A S L K Q P Y I T Q N Y
 F P V G T V V E Y E C R P G Y R R E P S L
 S P K L T C L Q N L K W S T A V E F C K K
 K S C P N P G E I R N G Q I D V P G G I L
 F G A T I S F S C N T G Y K L F G S T S S
 F C L I S G S S V Q W S D P L P E C R E I
 Y C P A P P Q I D N G I I Q G E R D H Y G
 Y R Q S V T Y A C N K G F T M I G E H S I
 Y C T V N N D E G E W S G P P P E C
 S S P N K C T P P N V E N G I L V S D N
 R S L F S L N E V V E F R C Q P G F V M
 K G P R R V K C Q A L N K W E P E L P S
 C S R V C Q P P P D V L H A E R T Q R D
 K D N F S P G Q E V F Y S C E P G Y D L
 R G A A S M R C T P Q G D W S P A A P T
 C E V K S C D D F M G Q L L N G R V L F
 P V N L Q L G A K V D F V C D E G F Q L
 K G S S A S Y C V L A G M E S L W N S S
 V P V C E Q I F C P S P P V I P N G R H
 T G K P L E V F P F G K A V N Y T C D P
 H P D R G T S F D L I G E S T I R C T S
 D P Q G N G V W S S P A P R C G I L G H
 C Q A P D H S F L F A K L K T Q T N A S D
 F P I G T S L K Y E C R P E Y Y G R P F
 S I T C L D N L V W S S P K D V C K R K
 S C K T P P D P V N G M V H V I T D I Q
 V G S R I N Y S C T T G H R L I G H S S
 A E C I L S G N A A H W S T K P P I C Q
 R I P C G L P P T I A N G D F I S T N R
 E N F H Y G S V V T Y R C N P G S G G R
 K V F E L V G E P S I Y C T S N D D Q V
 G I W S G P A P Q C I I P N K C T P P N

V E N G I L V S D N R S L F S L N E V V
 E F R C Q P G F V M K G P R R V K C Q A
 L N K W E P E L P S C S R V C Q P P P D
 V L H A E R T Q R D K D N F S P G Q E V
 F Y S C E P G Y D L R G A A S M R C T P
 Q G D W S P A A P T C E V K S C D D F M
 G Q L L N G R V L F P V N L Q L G A K V
 D F V C D E G F Q L K G S S A S Y C V L
 A G M E S L W N S S V P V C E Q I F C P
 S P P V I P N G R H T G K P L E V F P F
 G K A V N Y T C D P H P D R G T S F D L
 I G E S T I R C T S D P Q G N G V W S S
 P A P R C G I L G H C Q A P D H F L F A
 K L K T Q T N A S D F P I G T S L K Y E
 C R P E Y V G R P F S I T C L D N L V W
 S S P K D V C K R K S C K T P P D P V N
 G M V H V I T D I Q V
 G S R I N Y S C T T G H R L I G H S S
 A E C I L S G N A A H W S T K P P I C Q
 R I P C G L P P T I A N G D F I S T N R
 E N F H Y G S V V T Y R C N P G S G G R
 K V F E L V G E P S I Y C T S N D D Q V
 G I W S G P A P Q C I I P N K C T P P N
 V E N G I L V S D N R S L F S L N E V V
 E F R C Q P G F V M K G P R R V K C Q A
 L N K W E P E L P S C S R V C Q P P P D
 V L H A E R T Q R D K D N F S P G Q E V
 F Y S C E P G Y D L R G A A S M R C T P
 Q G D W S P A A P T C E V K S C D D F M
 G Q L L N G R V L F P V N L Q L G A K V
 D F V C D E G F Q L K G S S A S Y C V L
 A G M E S L W N S S V P V C E Q I F C P
 S P P V I P N G R H T G K P L E V F P F
 G K A V N Y T C D P H P D R G T S F D L
 I G E S T I R C T S D P Q G N G V W S S
 P A P R C G I H H H H H H H

Fig 9B

SEQ. ID NO:16

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 GACTGTGGCCTTCCCCCAGATGTACCTAATGCCAGCCAGCTTTGGAAGGCCGTACAAGTTTT
 CCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTTGTGAAAATTCTGGCGAGAAG
 GACTCAGTGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
 TGCGAGGTGCCAACAAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
 TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTTCTCTA

TCACCAAACTAACTTGCCCTTCAGAATTTAAATGGTCCACAGCAGTCGAATTTTGTA AAAAG
AAATCATGCCCTAATCCGGGAGAAATACGAAATGGTCAGATTGATGTACCAGGTGGCATATTA
TTTGGTGCAACCATCTCCTTCTCATGTAACACAGGGTACAAATTATTTGGCTCGACTTCTAGT
TTTTGTCTTATTTTCAGGCAGCTCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAGAAATT
TATTGTCCAGCACCACCACAAATTGACAATGGAATAATTCAAGGGGAACGTGACCATTATGGA
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TATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCCCACCACCTGAATGC
TCGAGTCCTAACAAATGCACGCCTCCAAATGTGGA AAAATGGAATATTGGTATCTGACAAC
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TGCTCCAGGGTATGTGAGCCACCTCCAGATGTCTGTCATGCTGAGCGTACCCAAAGGGAC
AAGGACAACCTTTTTCACCTGGGCAGGAAGTGTTCTACAGCTGTGAGCCCGGTACGACCTC
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TGTGAAGTGAAATCCTGTGATGACTTCATGGGCCAACTTCTTAATGGCCGTGTGCTATTT
CCAGTAAATCTCCAGCTTGGAGCAAAAGTGGATTTTGTGTTGTGATGAAGGATTTCAATTA
AAAGGCAGTCTGTGAAACAAATCTTTTGTCCAAGTCCTCCAGTTATTCTAATGGGAGACAC
ACAGGAAAACCTCTGGAAGTCTTTTCCCTTTGGAAAAGCAGTAAATTACACATGCGACCCC
CACCCAGACAGAGGGACGAGCTTCGACCTCATTGGAGAGAGCACCATCCGCTGCACAAGT
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TGTCAAGCCCAGATCATTTTCTGTTTGCCAAGTTGAAAACCCAAACCAATGCATCTGAC
TTTCCATTGGGACATCTTTAAAGTACGAATGCCGTCTGAGTACTACGGGAGGCCATTCT
TCTATCACATGTCTAGATAACCTGGTCTGGTCAAGTCCCAAAGATGTCTGTAAACGTAAA
TCATGTAAACTCCTCCAGATCCAGTGAATGGCATGGTGCATGTGATCACAGACATCCAG
GTGGATCCAGAATCAACTATTCTTGTACTACAGGGCACCGACTCATTGGTCACTCATCT
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CGAATTCCTTGTGGGCTACCCCCACCATCGCCAATGGAGATTTTATTAGCACCAACAGA
GAGAATTTTCACTATGGATCAGTGGTGACCTACCGCTGCAATCCTGGAAGCGGAGGGAGA
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GGCATCTGGAGCGGCCCGCCCCCTCAGTGCATTATACCTAACAAATGCACGCCTCCAAAT
GTGGAAAATGGAATATTGGTATCTGACAACAGAAGCTTATTTTCTTAAATGAAGTTGTG
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GTCTGTCATGCTGAGCGTACCCAAAGGGACAAGGACAACCTTTTCAACCGGGCAGGAAGTG
TTCTACAGCTGTGAGCCCGCTATGACCTCAGAGGGGCTGCGTCTATGCGCTGCACACCC
CAGGGAGACTGGAGCCCTGCAGCCCCCACATGTGAAGTGAAATCCTGTGATGACTTCATG
GGCCAACCTTCTTAATGGCCGTGTGCTATTTCCAGTAAATCTCCAGCTTGGAGCAAAAGTG
GATTTTGTGTTGTGATGAAGGATTTCAATTAAGGAGCAGCTCTGCTAGTTATTGTGTCTTG
GCTGGAATGGAAAACCTTTGGAATAGCAGTGTTCCAGTGTGTGAACAAATCTTTTGTCCA
AGTCTCCAGTTATTCTTAATGGGAGACACACAGGAAAACCTCTGGAAGTCTTTCCCTTT
GGAAAAGCAGTAAATTACACATGCGACCCCCACCCAGACAGAGGGACGAGCTTCGACCTC
ATTGGAGAGAGCACCATCCGCTGCACAAGTGACCTCAAGGGAATGGGGTTTGGAGCAGC
CCTGCCCCCTCGCTGTGGAATCTGGGTCACTGTCAAGCCCCAGATCATTTTCTGTTTGCC
AAGTTGAAAACCCAAACCAATGCATCTGACTTTCCCATTTGGGACATCTTTAAAGTACGAA
TGCCGTCTGAGTACTACGGGAGGCCATTCTCTATCACATGTCTAGATAACCTGGTCTGG
TCAAGTCCCAAAGATGTCTGTAAACGTAAATCATGTAAACTCCTCCAGATCCAGTGAAT
GGCATGGTGCATGTGATCACAGACATCCAGGTT
GGATCCAGAATCAACTATTCTTGTACTACAGGGCACCGACTCATTGGTCACTCATCT
GCTGAATGTATCCTCTCGGGCAATGCTGCCCATTGGAGCACGAAGCCGCCAATTTGTCAA
CGAATTCCTTGTGGGCTACCCCCACCATCGCCAATGGAGATTTTATTAGCACCAACAGA
GAGAATTTTCACTATGGATCAGTGGTGACCTACCGCTGCAATCCTGGAAGCGGAGGGAGA

R S L F S L N E V V E F R C Q P G F V M
 K G P R R V K C Q A L N K W E P E L P S
 C S R V C Q P P P D V L H A E R T Q R D
 K D N F S P G Q E V F Y S C E P G Y D L
 R G A A S M R C T P Q G D W S P A A P T
 C E V K S C D D F M G Q L L N G R V L F
 P V N L Q L G A K V D F V C D E G F Q L
 K G S S A S Y C V L A G M E S L W N S S
 V P V C E Q I F C P S P P V I P N G R H
 T G K P L E V F P F G K A V N Y T C D P
 H P D R G T S F D L I G E S T I R C T S
 D P Q G N G V W S S P A P R C G I L
 V E S K Y G P P C P S C P A P E F L
 G G P S V F L F P P K P K D T L M I S R
 T P E V T C V V V D V S Q E D P E V Q F
 N W Y V D G V E V H N A K T K P R E E Q
 F N S T Y R V V S V L T V L H Q D W L N
 G K E Y K C K V S N K G L P S S I E K T
 I S K A K G Q P R E P Q V Y T L P P S Q
 E E M T K N Q V S L T C L V K G F Y P S
 D I A V E W E S N G Q P E D N Y K T T P
 P V L D S D G S F F L Y S R L T V D K S
 R W Q E G N V F S C S V M H E A L H N H
 Y T Q K S L S L S P G K

Fig. 11B
SEQ. ID NO:20

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 CCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTTGTGAAAATTCTGGCGAGAAG
 GACTCAGTGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
 TGCGAGGTGCCAACAAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
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 AAAGGACCCCGCGGTGTGAAGTGCCAGGCCCTGAACAAATGGGAGCCGGAGCTACCAAGC
 TGCTCCAGGGTATGTGAGCCACCTCCAGATGTCTGTCATGCTGAGCGTACCCAAAGGGAC
 AAGGACAACTTTTACCTGGGCAGGAAGTGTCTACAGCTGTGAGCCCGGCTACGACCTC

ATA TAC GAA TTC TGG GTC ACT GTG AGG AGC CAC CAA CAT TTG AAG C

ACA GTG AGA TCT TTA GTG ATG GTG ATG GTG ATG CGA CAC TTT AAG ACA CTT
TGG AAC

[illegible]

SEO. ID NO:24

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CTCCTCGGGGAGCTGCCCGGCTGCTGCTGCTGGTGTGTTGTGCCCTGCCGGCCGTGTGGGGT
GACTGTGGCCCTTCCCCCAGATGTACCTAATGCCCAGCCAGCTTTTGGGAAGGCCGTACAAGTTT
CCCAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTTGTGAAAATTCTTGCGCAGAAG
GACTCAGTGATCTGCCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
TGCGAGGTGCCAACAAAGGCTAAATTCTGCATCCCTCAAACAGCCCTTATATCACTCAGAATTAT
TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTTCTCTA
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ACAGGAAAACCTCTGGAAGTCTTTCCCTTTGGAAAAGCAGTAAATTACACATGCGACCCC
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TGTGAGGAGCCACCAACATTTGAAGCTATGGAGCTCATTTGGTAAACCAAAACCCCTACTAT
GAGATTGGTGAACGAGTAGATTATAAGTGTAAGGATACTTCTATATACCTCCTCTT
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GGTGAAGAAATTCATATTTGTGAACTTAAAGGATCAGTAGCAATTTGGAGCGGTAAGCCC
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CCTGGACCAGATCCATTTTCACTTATTGGAGAGAGCACGATTTATTGTGGTGACAATTCA
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GAATGCGATAAGGGTTTTTACCTCGATGGCAGCGACACAATTGTCTGTGACAGTAACAGT
ACTTGGGATCCCCAGTTCCAAAGTGCTTAAA//GTGTCG//CATCACCATCACCATCAC
TAAAGATCT

WESTERN BLOT OF HYBRID PROTEINS DAF-IgG4, DAF-CR1BB, and DAF-CR1B

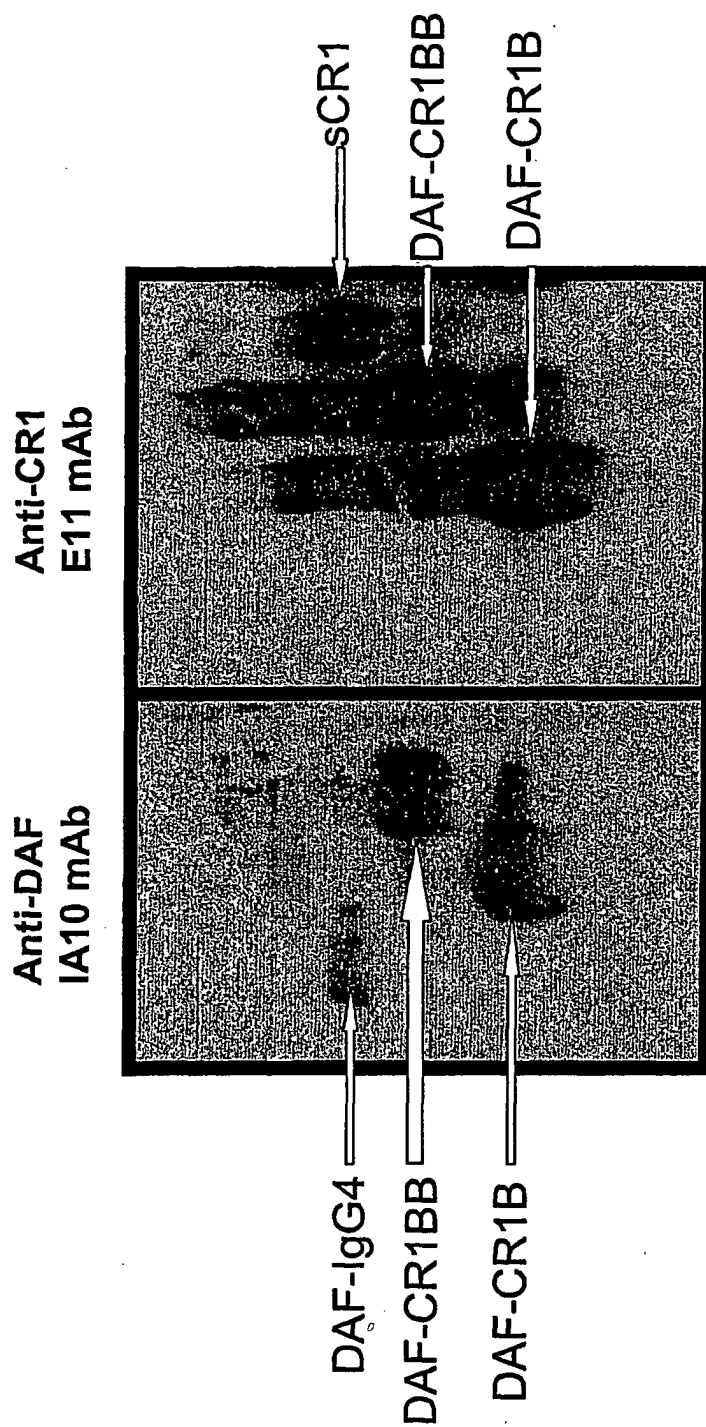


Fig. 14

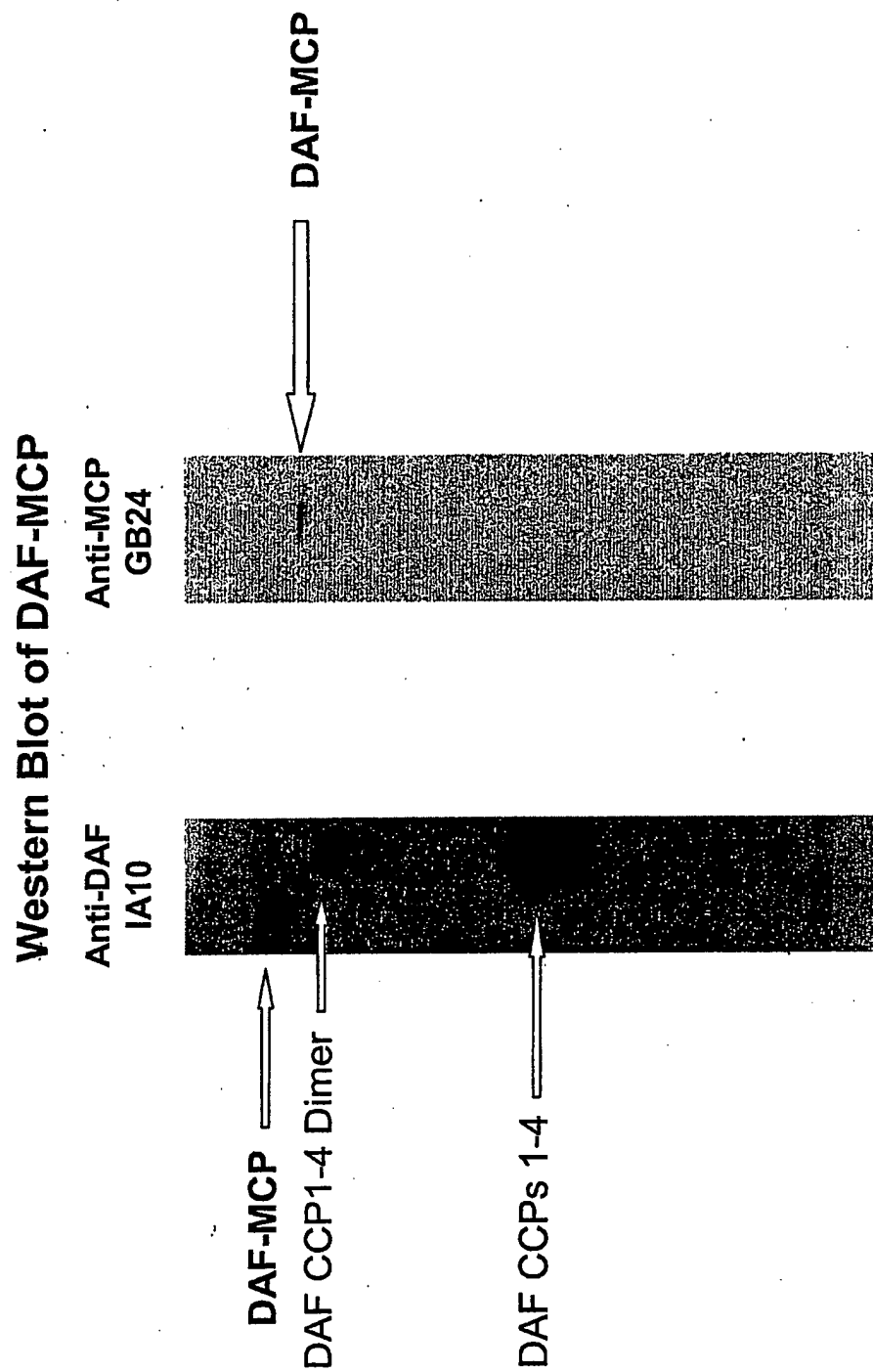


Fig. 15

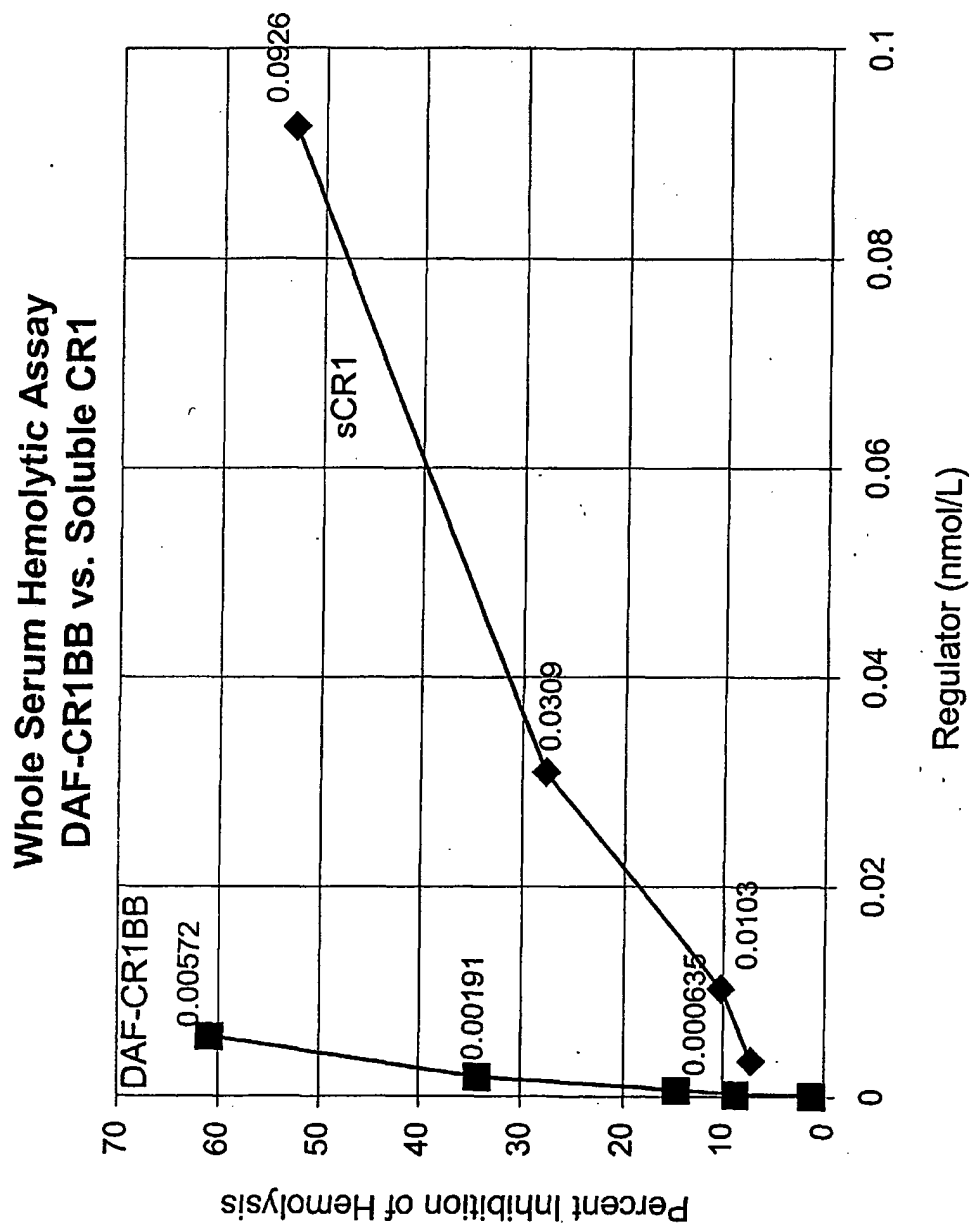


Fig. 16

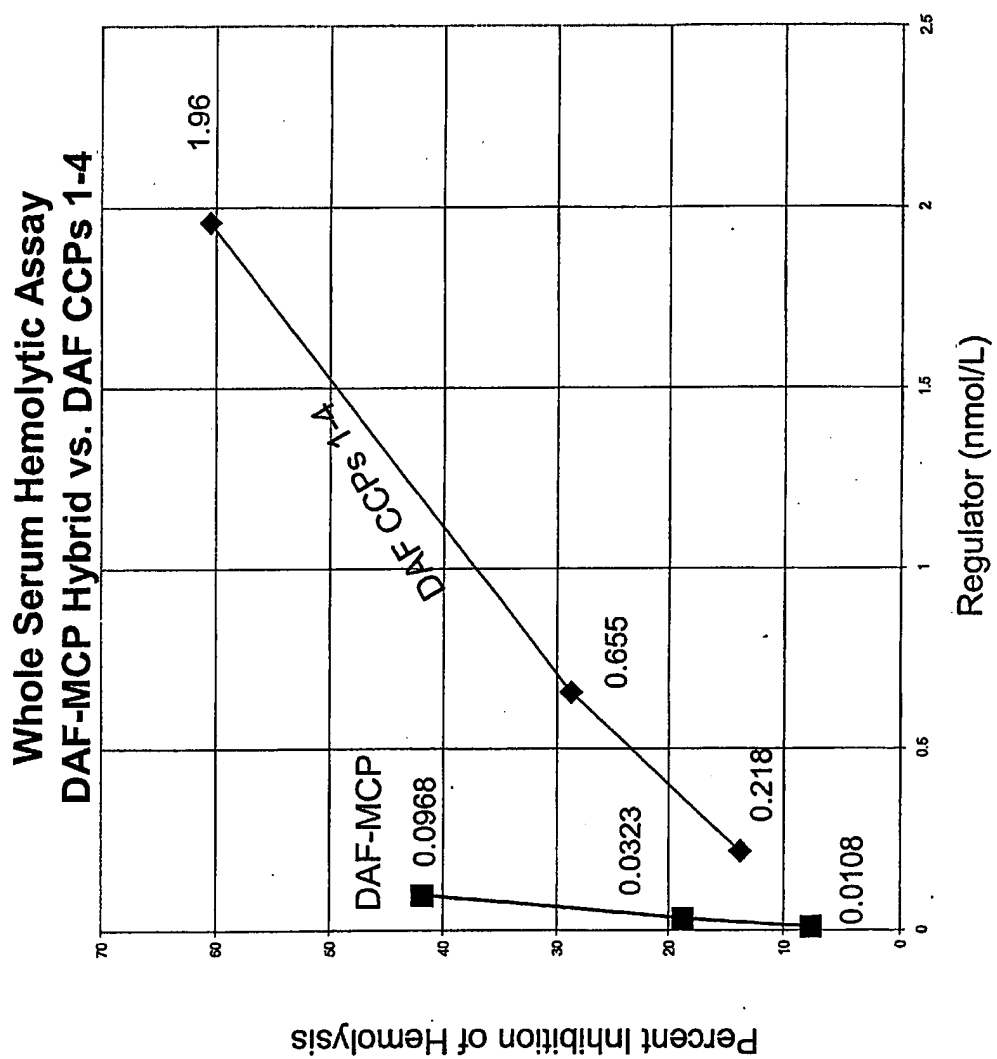


Fig. 17

Classical Pathway C3 Convertase Decay

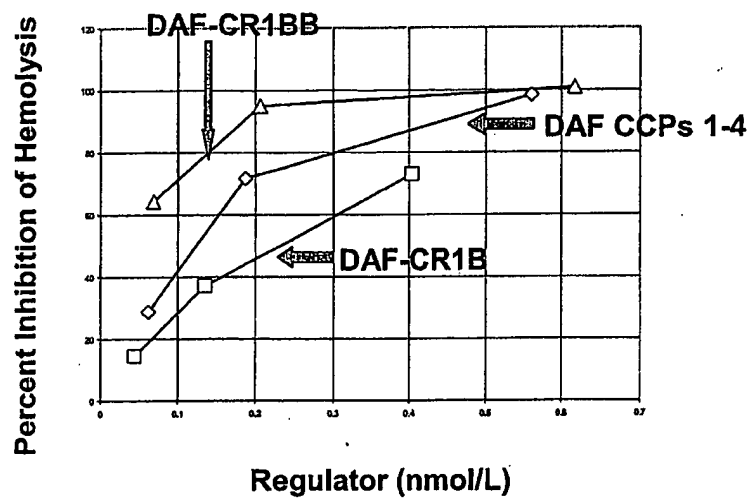


Fig. 18A

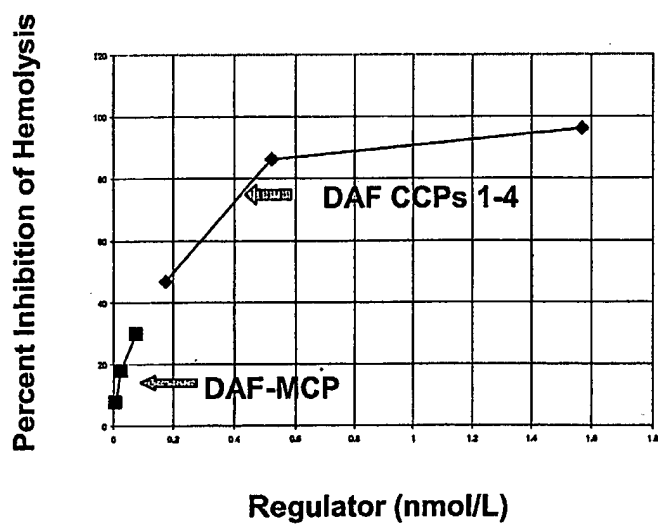


Fig. 18B

Classical Pathway C5 Convertase Decay DAF-CR1B vs DAF CCPs 1-4

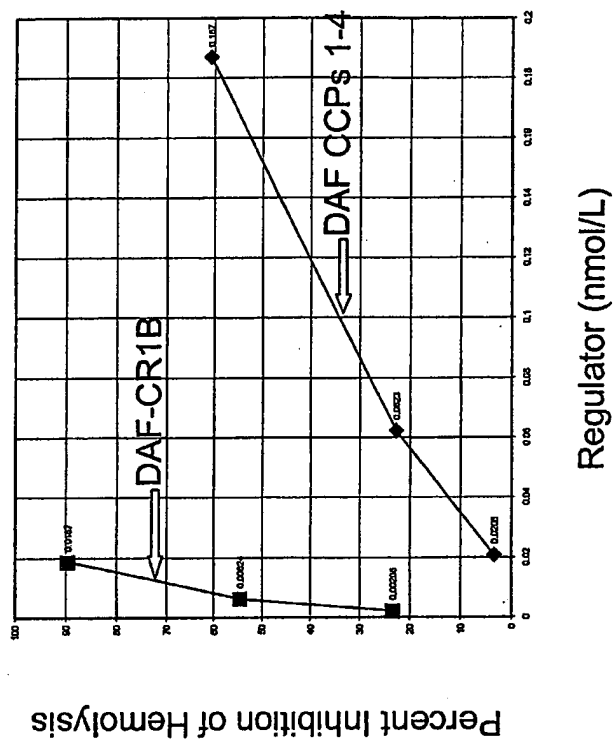


Fig. 19

Classical Pathway C5 Convertase Decay DAF-CR1BB vs sCR1 vs DAF-CR1B

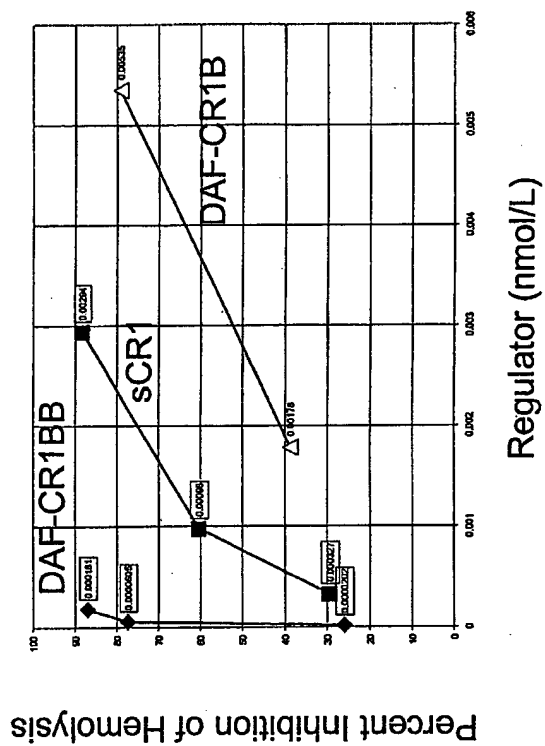


Fig. 20

Cell-bound (E^{sh}C4b3b) Cofactor Assay

Cell Supernatant

(Anti-human C3 pAb)

COS SN DAF-MCP DAF-CR1BB

+I +I +I

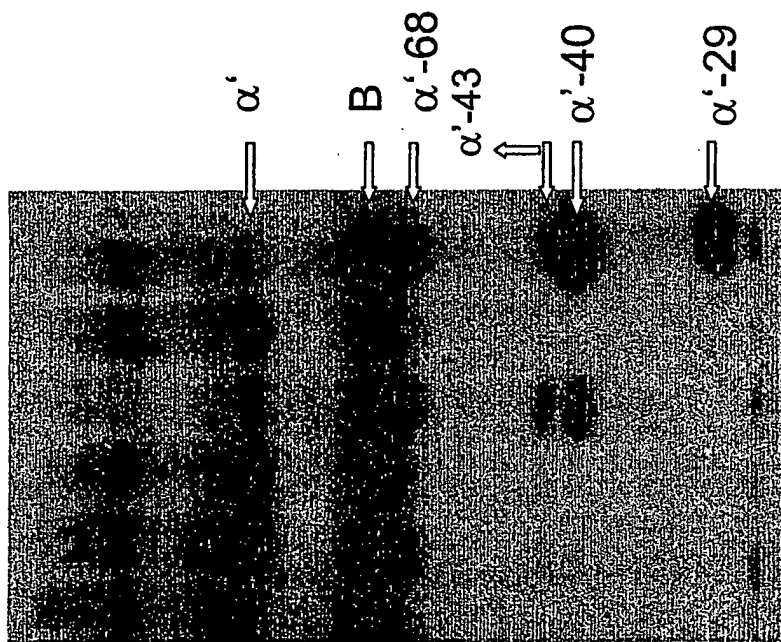


Fig. 21

Cell-bound (EshC4b3b) Cofactor Assays Cell Supernatant

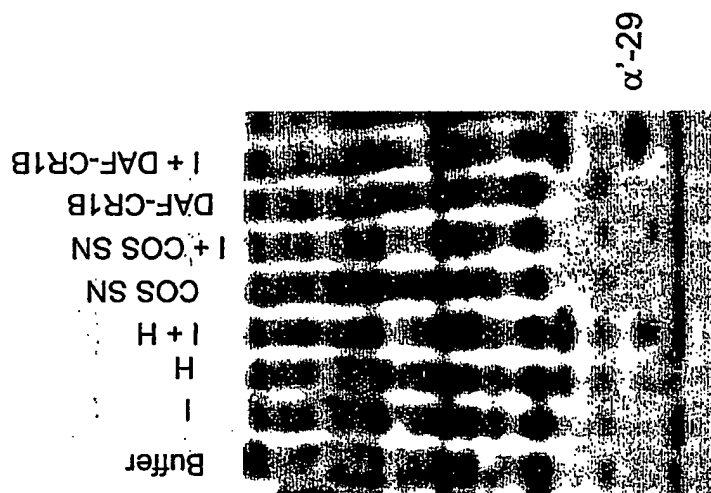


Fig. 22